

TCG ACC CAC GCG TCC GGC CTT GGC AGA GTC TGG GGT CCC TGG ACT GAG CCA TCA GCT GGG  
TCA CTG AGA CCC ATG GCA AGG AAA CAA AAT AGG AAT TCC AAG GAA CTG GGC CTA GTT CCC

CTC ACA GAT GAC ACC AGC CAC GCC GGG CCT CCA GGG CCA GGG AGG GCA CTG CTG GAG TGT  
GAC CAC CTG AGG AGT GGG GTG CCA GGT GGA AGG AGA AAG GAC TGG TCC TGC TCG CTC

CTC GTG GCC TCC CTC GCG GGC GCC TTC GCC TCC ATC AAG GCC TTT TAC AAT GAG TCA TGG GAA AGA AGG  
GTG GTG AAT GCC CCC ACC CCG TAC ATC AAG GCC TTT TAC AAC TAC GGC TAC AAC CTG TCG  
CAT GGA CGT CCA ATA GAC CCA GAC ACT CTG ACT CTG CTC TGG TGT GTG ACT GTG TCC ATG

TTC GCC ATC GGT GGA CTT GTG GGG ACG TTA ATT GTG AAG ATT GGA AAG GTT CTT GGG

**FIG. 1A**

AGG AAG CAC ACT TTG CTG GCC AAT AAT GGG TTT GCA ATT TCT GCT GCA TTG CTG ATG GCC  
TGC TCG CTC CAG GCA GGA GCC TTT GAA ATG CTC ATT GTG GGA CGC TTC ATC ATG GGC ATA  
GAT GGA GGC GTC GCC CTC AGT GTG CTC CCC ATG TAC CTC AGT GAG ATC TCA CCC AAG GAG  
ATC CGT GGC TCT CTG GGG CAG GTG ACT GCC ATC TTT ATC TGC ATT GGC GTG TTC ACT GGG  
CAG CTT CTG GGC CTG CCC GAG CTG CTG GGA AAG GAG AGT ACC TGG CCA TAC CTG TTT GGA  
GTG ATT GTG GTC CCT GCC GTT GTC CAG CTG CTG AGC CTT CCC TTT CTC CCG GAC AGC CCA  
CGC TAC CTG CTC TTG GAG AAC CAC AAC GAG GCA AGA GCT GTG AAA GCC TTC CAA ACG TTC

**FIG. 1B**

TTG GGT AAA GCA GAC GTC TCC CAA GAG GTA GAG GAG GTC CTG GCT GAG AGC CAC GTG CAG  
AGG AGC ATC CGC CTG GTG TCC GTG CTG GAG CTG CTG AGA GCT CCC TAC GTC CGC CGC TGG CAG  
GTG GTC ACC GTG ATT GTC ACC ATG GCC ATG GCC TGC TAC CAG CTC TGT GGC CTC AAT GCA ATT TGG  
TTC TAT ACC AAC AGC ATC TTT GGA AAA GCT GGG ATC CCT CCG GCA AAG ATC CCA TAC GTC  
ACC TTG AGT ACA GGG GGC ATC GAG ACT TTG GCT GCC GTC TTC TCT GGT TTG GTC ATT GAG  
CAC CTG GGA CGG AGA CCC CTC CTC ATT GGT GGC TTT GGG CTC ATG GCC CTC TTC TTT GGG  
ACC CTC ACC ATC ACG CTG ACC CTG CAG GAC CAC GCC CCC TGG GTC CCC TAC CTG AGT ATC  
GTG GGC ATT CTG GCC ATC ATC GCC TCT TTC TGC AGT GGG CCA GGT GGC ATC CCG TTC ATC

**FIG. 1C**

TTG ACT GGT GAG TTC TTC CAG CAA TCT CAG CCG CCG GCT GCC TTC ATC ATT GCA GGC ACC  
GTC AAC TGG CTC TCC AAC TAC TTT GCT GGT GTT GGG CTC CTC TTC CCA TTC ATT CAG AAA AGT CTG  
GAC ACC TAC TGT TTC CTA GTC TTT GCT ACA ATT TGT ATC ACA GGT GCT ATC TAC CTG TAT  
TTT GTG CCT GAG ACC AAA AAC AGA ACC TAT GCA GAA ATC AGC CAG GCA TTT TCC AAA  
AGG AAC AAA GCA TAC CCA CCA GAA GAG AAA ATC GAC TCA GCT GTC ACT GAT GCT CCT GCT  
TCT CCT ACT ACT CCG AAT ACA GCC TGG ATT CAA GCT GCC GCC ACC ACC ACC ACC GCC  
ACC AAA GAA CAC CCA TTG TAA ACG GTC ATG TGG TAT TTC CTC AAC CTC GAA TGA CCT

**FIG. 1D**

TCC	CCT	ATC	TTC	TCC	TGG	AGA	ACA	CCA	AGT	CAT	GAT	GTC	AGA	CAA	GAG	CTT	GGA	TTT
TGG	AGA	CAT	GGG	TTT	GAA	TTC	CAG	TCA	TTC	ATT	CTT	TTA	AGC	AAA	TAT	TTA	ACA	AGT
ACT	GAC	ATG	TCC	CAT	ATG	TTG	TTC	TAC	CCA	CTG	GTT	ATA	CAA	TGG	GAG	GGA	GAG	GAG
AGA	GAG	AGA	GAG	GAT	GCT	ATT	CTA	AAA	GCT	TGA	AGT	CTA	GGC	TGT	GCA	CGG	TGG	CTC
ACG	CCT	GTA	ATC	CCA	GCA	CTT	TGG	GAG	GCC	GAG	GTG	GGT	GGA	TGA	GGT	CAG	GAG	ATT
GAG	ACC	ATC	CTG	GCT	AAC	ATG	GTG	AAA	CTC	CCT	TAC	TAA	AAA	TAC	AAA	AAA	TTA	GCT
GAG	CAT	GGT	GGC	GGG	CGC	CTG	TAG	TCC	CAG	CTA	CTT	GGG	AGG	CTG	AGG	CAG	GAG	AAT
GTG	AAC	CCA	GGA	GGA	GCT	TGC	AGT	GAG	CCG	AGA	TCA	CAC	CAC	ACT	CCA	GCC	TGG	GGC
GTG	ACA	GAG	CCA	GAC	TCC	GTC	TCA	AAA	AAG	GGC	GGC	GGC						

FIGURE

met ala arg lys gln asn arg asn ser lys glu leu alv alv leu val pro  
leu thr asp thr ser his ala gly pro pro gly pro gly arg ala leu leu glu cys  
asp his leu arg ser gly val pro gly gly arg arg lys asp trp ser cys ser leu  
leu val ala ser leu ala gly ser ser phe leu tyr gly tyr asn leu ser  
val val asn ala pro thr pro tyr ile lys ala phe tyr asn glu ser trp glu arg arg  
his gly arg pro ile asp pro asp thr leu thr leu leu trp ser val thr val ser ile  
phe ala ile gly gly leu val gly thr leu ile val lys met ile gly lys val leu gly

**FIG. 2A**

arg lys his thr leu leu ala asn asn gly phe ala ile ser ala ala leu leu met ala  
cys ser leu gln ala gly ala phe glu met leu ile val gly arg phe ile met gly ile  
asp gly val ala leu ser val leu pro met tyr leu ser glu ile ser pro lys glu  
ile arg gly ser leu gly gln val thr ala ile phe ile cys ile gly val phe thr gly  
gln leu gly leu pro glu leu leu gly lys glu ser thr trp pro tyr leu phe gly  
val ile val val pro ala val val gln leu leu ser leu pro phe leu pro asp ser pro  
arg tyr leu leu glu lys his asn glu ala arg ala val lys ala phe gln thr phe

**FIG. 2B**

leu gly lys ala asp val ser gln glu val glu val leu ala glu ser his val gln  
arg ser ile arg leu val ser val leu glu leu leu arg ala pro tyr val arg trp ala  
val val thr val ile val thr met ala cys tyr gln leu cys gly leu asn ala ile trp  
phe tyr thr asn ser ile phe gly lys ala gly ile pro pro ala lys ile pro tyr val  
thr leu ser thr gly gly ile glu thr leu ala ala val phe ser gly leu val ile glu  
his leu gly arg pro leu leu ile gly gly leu met gly leu phe phe gly  
thr leu thr ile thr leu thr leu gln asp his ala pro trp val pro tyr leu ser ile  
val gly ile leu ala ile ile ala ser phe cys ser gly pro gly gly ile pro phe ile

**FIG. 2C**

Leu thr gly glu phe phe gln gln ser gln arg pro ala ala phe ile ile ala gly thr  
val asn trp leu ser asn phe ala val gly leu leu phe pro phe ile gln lys ser leu  
asp thr tyr cys phe leu val phe ala thr ile cys ile thr gly ala ile tyr leu tyr  
phe val leu pro glu thr lys asn arg thr tyr ala glu ile ser gln ala phe ser lys  
arg asn lys ala tyr pro pro glu glu lys ile asp ser ala val thr asp ala pro ala  
ser ser pro phe thr thr pro asn thr ala trp ile gln ala ala ala thr thr thr ala  
thr lys lys glu his pro leu

**FIG. 2D**

Figure 3A of 4

**FIG. 3A**

MLVNNVLAIAAGGLEMGGLAKXAXSFEMLILG Majority

190 200 210

MLVNNVLSIAGNLLMGGLAKMGPHILIAAG glut1  
 MLIVNLLAIAAGGCLMGFCKIAESVEMLILG glut2  
 MLVNNVIAVLAGGSLMGLANAAASYEMLILG glut3  
 LIFNNIFSIVPAILMGCSRVATTFELIIS glut4  
 LANNGFAISALLMACSLQAGAEEMLITVG glut5  
 LANNGFAISALLMACSLQAGAEEMLITVG GlutX

RFIIGLYCGLSSGVVPMYVGEISPTALRGA Majority

220 230 240

RFIIGVYCGLTTTGFVPMYVGEVCPTEELRGA glut1  
 RAITGLYCGLSSGLVPMYVSEVECPTALRGA glut2  
 RLTIGLFCGLCTGFVPMYIGEGEISPTALRGA glut3  
 RFLIGLGAYSTSGLGLVPMYVGETAPTHLRGA glut4  
 RLLVGICAGVVSSNVVPMYLCEAEKNILRGA glut5  
 RFIMGIDGGVALSVLPMYLSETSPKEIPGS GlutX

LGTLNOLGIVIGILIAQVLGLDSLLGNESL Majority

250 260 270

LGTLHQLGIVVGGIILIAQVFGLDSIMGNQEL glut1  
 LGTLHQLAIVTGILISQVLGDFILGNDEE glut2  
 FGTLNOLGIVIGILVAQIFGLKVIGTEDL glut3  
 LGTLNOLAIVITGILIAQVLGLEESLLGTASL glut4  
 LGVVPOLFITVGILVAQIFGLRNLLANVDG glut5  
 LGQVTAIFCIGVFTGOLGLPELLGKEST GlutX

WPLLGLTGVPALLLLLPFCPESPRYLL Majority

280 290 300

WPLLSVIFPALLQCIELPFCPESPRFLL glut1  
 WPLLGISGVAALLQFFLLLCPESPRYY glut2  
 WPLLGFTIPAIIQCAALPFFCPESPRFLL glut3  
 WPLLGLTVLPALLQLVLLPFCPESPRYY glut4  
 WPLLGLTGVPAALQLLLPFFPESPRYY glut5  
 WPYIFGVIVPAVVOLLSIPFLPDSPRYLL GlutX

INKNEEARAKKALQRLRGTADVSQEEVAEMK Majority

310 320 330

INRNEENNRAKSVLKLRGTADVTRDLQEMK glut1  
 IKLGKVEEAKKSLKILRGNCDPMKEIAEME glut2  
 INRKEEEEKAKEILQRLWGTEDVAQDIQEMK glut3  
 IIQNLEGPARKSLKITGWADVSGVLAELK glut4  
 IIQKKDEAAAKKAQTIRGWDSVDREVAEIR glut5  
 LEKHNEARAVKAFQTFLGKADVSOEVEEVL GlutX

DESRXMXSEKXVSLEFRSRXYRQPVVIA Majority

340 350 360

EEESRQMMREKKVTIELFRSAAYRQPILIA glut1  
 KEKQEAASEKRVSIGQFSSSKYRQAVIVA glut2  
 DESMRMSQEKQVTIELFRAPNYRQPIIS glut3  
 DEKRKLEREPLSLQLLGSRTHRQPLITA glut4  
 QEDEAEKAAGFISVLKLFRMRSLWQLSI glut5  
 AFSHVQRSIRLVVSIELLRAPYVRWQVTV GlutX

IVLQLSQQLSGINAVFYYSSTSIEEKAGVGO Majority

370 380 390

167 VVLQLSQQLSGINAVFYYSSTSIEEKAGVQQ  
320 LMVQIISQQFSGINAIIFYYSTNIEQRAAGVGO  
274 IMLQLSQQLSGINAVFYYSSTSIEEKAGVQE  
292 VVLQLSQQLSGINAVFYYSSTSIEEKAGVGO  
282 IVLMMGGOQLSGVNAIYVYADQIYLSAGVPE  
321 IVTMACYOLCGLMAIWFWYTNITFGRAGIIPP  
giut1-pro  
giut2-pro  
giut3-pro  
giut4-pro  
giut5-pro  
GlutX-pro  
final-2/6/98

P - - VYATIGAVVNTVFTVVSVFVVERAGR Majority

400 410 420

197 P - - VYATIGSGIVNTAFTVVVCLFVVERAGR  
350 P - - VYATIGVGVVNTVFTVIVSVFLVERAGR  
304 P - - VYATIGAGVVNTIPTVVVSVFLVERAGR  
322 P - - AYATIGAGVVNTVFTLVSVLLVERAGR  
312 EH VQ VTAGTGA VNVVMTFCAVFVVELLGR  
351 AKI PYVTLSTGIE TLA AVFSGLWIFHLGR  
giut1-pro  
giut2-pro  
giut3-pro  
giut4-pro  
giut5-pro  
GlutX-pro  
final-2/6/98

RTLHLGLGGMAGCAVLMTIALALLDQVPW Majority

430 440 450

225 RTLHLIGLAGMAGCAVLMTIALALLEQOLPW  
378 RSLFЛАGLMGM LISAVAMTVGVVILSQAFAW  
332 RTLHLIGLGGMAFCISIMMTISLLKDNYSW  
350 RTLHLLLCGLAGMCGCAILMTVALLLERVPW  
342 RLLL LLLGFSICLIAACCVLTAALALQDTVSW  
381 RPLLIGGFGLMGLFFGTLTITETLQDHAPW  
giut1-pro  
giut2-pro  
giut3-pro  
giut4-pro  
giut5-pro  
GlutX-pro  
final-2/6/98

MSYVSIVAIFFGVAFFEVGPGPPIPWFIVAE Majority

460 470 480

255 MSYLSIVAIFFGVAFFEVGPGPPIPWFIVAE  
408 MSYVSMVAIFLFVIVFEEVGPGPPIPWFIVAE  
362 MSFICIGAIILVVFVAFFEIIGPGPPIPWFIVAE  
380 MSYVSVIVAIFFGVAFFEIIGPGPPIPWFIVAE  
372 MPYISIVCVISYVIGHALGPSPIPALLITE  
411 VPYLSIVGILAIIASFCSGPGGTIPFILTGE  
giut1-pro  
giut2-pro  
giut3-pro  
giut4-pro  
giut5-pro  
GlutX-pro  
final-2/6/98

LFSQGPRAAIAVAGFSNWTSNFIVGLLFQ Majority

490 500 510

285 LFSQGPRAAIAVAGFSNWTSNFIVGMCFO  
438 LFSQGPRAAIAVAGFCNWACNFTVGMCCFO  
392 LFGQGPRAAMAVAGCNSWTSNFLVGYLIFP  
410 LFSQGPRAAMAVAGFSNWTSNFIICMGFO  
402 IFLQSSRPSAFMVGGSVHWLSNFTVGLIFP  
441 FFOQOSQRPAAFITAGTVWLNFAVGILLF  
giut1-pro  
giut2-pro  
giut3-pro  
giut4-pro  
giut5-pro  
GlutX-pro  
final-2/6/98

YIAELLGPYVFIVFAVLLLFIFTFLKVP Majority

520 530 540

315 YVEQLCGPYVFTIFTVLLVLFIFTYFKVP  
468 YIADLCGPYVFFVVFVAVLLLVFELFAYIKVP  
422 SATFYLGAYVFTIVFTVFLVIFWVFTFFKVP  
440 YVAEAMGPYVFLLFVAVLLLGFFIFTFLRVP  
432 FIQEGLGPYSFTIVFAVIVCLITIYIFLIVP  
471 FTQKSIDTYCFLVFAATICITGAIYLYFVLP  
giut1-pro  
giut2-pro  
giut3-pro  
giut4-pro  
giut5-pro  
GlutX-pro  
final-2/6/98

Majority

E T K G R T F D E I A A A F R K X N K X - E Q - P E K E S I	Majority	
550	560	570
E T K G R T F D E I A S G F R Q G G A - - S Q - S D E T P E	glut1	
E T K G K S F E E I A A A F R R R K K L P A K - - - - - M	glut2	
E T R G R T F E E I T R A F E G Q V Q T G T R - G E K G P I	glut3	
E T R G R T F D Q I S A A F H R T P S L L E O - E V K P S T	glut4	
E T K A K T F I E I N Q I F T K M N K V S E V Y P E K E E L	glut5	
E T K N R T Y A E I S Q A F S K R N K A - - - Y P P E E K T	GlutX	

Majority

E E L E P L G P D -	Majority	
580	590	600
F L F H P L G A D -	glut1	
T E E L E D I R G G -	glut2	
M E M N S I Q P T -	glut3	
E L - E Y L G P D -	glut4	
K E L P P V T S E -	glut5	
D S A V T D A P A S S P F T T P N T A W I Q A A A T T T A T	GlutX	

Majority

- - E - - X -	Majority
- - S Q V .	glut1
- - E - - E A	glut2
- - K D T N A	glut3
- - E N - D	glut4
- - - - - Q	glut5
K K E H P L .	GlutX

FIG. 3D

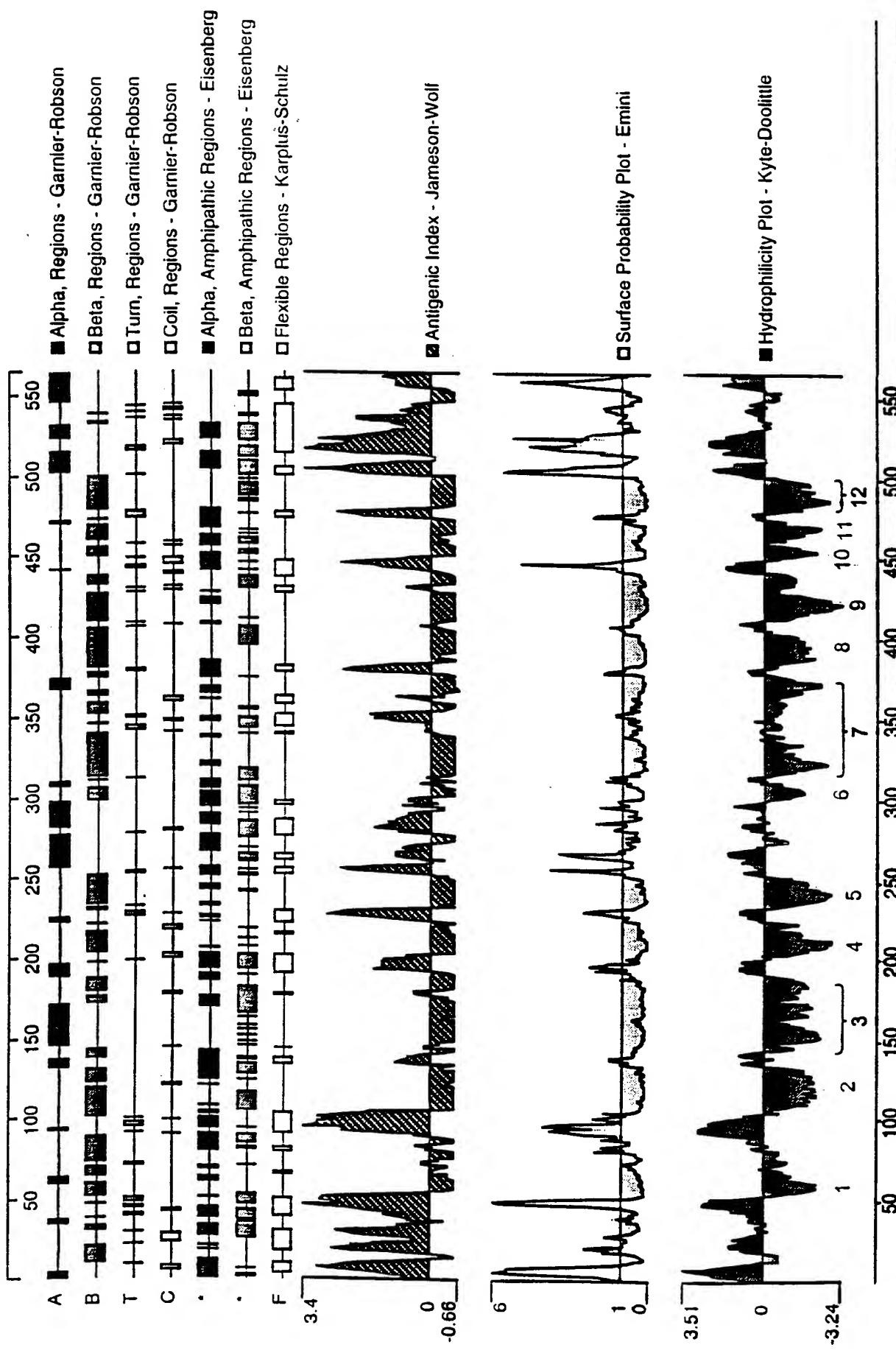


FIG. 4